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2 Moderate alcohol exposure during early brain development increases
3 stimulus-response habits in adulthood

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22 **Abstract**

23 Exposure to alcohol during early central nervous system (CNS) development has
24 been shown variously to affect aspects of physiological and behavioural
25 development. In extreme cases, this can extend to craniofacial defects, severe
26 developmental delay and mental retardation. At more moderate levels, subtle
27 differences in brain morphology and behaviour have been observed. One clear
28 effect of developmental alcohol exposure is an increase in the propensity to
29 develop alcoholism and other addictions. The mechanisms by which this occurs,
30 however, are not currently understood. In this study we tested the hypothesis that
31 adult zebrafish chronically exposed to moderate levels of ethanol during early
32 brain ontogenesis would show an increase in conditioned place preference (CPP)
33 for alcohol and an increased propensity towards habit formation, a key
34 component of drug addiction in humans. We found support for both of these
35 hypotheses and found that the exposed fish had changes in mRNA expression
36 patterns for dopamine receptor, nicotinic acetylcholine receptor (nAChR) and μ -
37 opioid receptor (MOR) encoding genes. **Collectively, these data show an**
38 **explicit link between the increased proclivity for addiction and addiction-**
39 **related behaviour following exposure to ethanol during early brain**
40 **development, and alterations in the neural circuits underlying habit**
41 **learning.**

42

43

44 **Keywords:** CPP; moderate prenatal ethanol; habit; zebrafish; addiction; μ -opioid;
45 dopamine

46 Introduction

47 Maternal alcohol consumption during pregnancy results in a range of effects on
48 the developing foetus, collectively referred to as Foetal Alcohol Spectrum
49 Disorders (FASD; Paintner et al., 2012). FASD are characterized by a range of
50 teratogenic and psychological defects, and represent the leading non-hereditary
51 cause of mental retardation, with the prevalence estimated at between 2 and 5%
52 of the population of the USA and western Europe (May et al., 2009). In extreme
53 circumstances, when mothers drink heavily during pregnancy, foetal alcohol
54 syndrome (FAS) results in gross skeletal and craniofacial abnormalities, and
55 severe CNS dysfunction (Hanson et al., 1976). More moderate alcohol
56 consumption (e.g., equivalent to 1-2 drinks/day, average BAC ~0.01-0.04g/dL
57 Valenzuela et al., 2012), however, is associated with a range of subtle cognitive
58 and behavioural defects, including aggression and affective disorders (Sood et
59 al., 2001), impulsivity and inattention (Streissguth et al., 1989; Suess et al., 1997),
60 poor scholastic performance (Olson et al., 1998) and deficits in social behaviour
61 (Keil et al., 2010; Kelly et al., 2000; Thomas et al., 1998).

62 Exposure to moderate ethanol during gestation also poses an increased
63 risk that the offspring will develop substance abuse problems in later life
64 (Valenzuela et al., 2012). For example, Baer et al. (2003) demonstrated in a 21-
65 year prospective study that adults who had been exposed to ethanol during
66 gestation had a higher incidence of alcohol abuse. This study, and others like it
67 (Alati et al., 2006; Alati et al., 2008; Miller, 1999; Streissguth et al., 1994) have
68 typically controlled very robustly for the effects of post-natal extraneous
69 environmental mediators (e.g., parental role models, etc.), and there is also
70 evidence from adoption studies that suggest direct cause-effect relationships

71 (Landgren et al., 2010; van Wieringen et al., 2010). Nevertheless, clinical studies
72 have only limited efficacy in describing the cellular and molecular mechanisms by
73 which ethanol exposure during gestation exerts its effects on addiction in later life.

74 Preclinical studies using animal models have proved very useful in this
75 regard for a number of reasons. First, it is possible to titrate alcohol intake in a
76 precise manner. Second, it is possible to exclude potential confounds (e.g., post-
77 natal environment, poly-drug abuse, etc.). Third, the gestational period during
78 which alcohol is present can be carefully scrutinised. Fourth, the offspring can be
79 monitored and tested at various developmental stages. For example, extensive
80 work with rodent models have variously demonstrated that ethanol during early
81 brain development affects key components of the mesoaccumbens dopamine
82 reward pathway, potentially explaining why affected individuals show greater drug
83 preference in later life (Blanchard et al., 1993; Fulginiti et al., 1994; Middaugh et
84 al., 1994; Wang et al., 2006).

85 Although these studies have been helpful in determining the impact of
86 developmental ethanol exposure on structural and functional alterations in
87 midbrain dopamine systems, simply increasing the reward potential of a drug is
88 not sufficient to explain the development of addiction to that (or other) drugs
89 (Everitt and Robbins, 2005; Robbins and Everitt, 1999). A key aspect of addiction
90 is the progressive 'loss of control' over drug use and the establishment of learned
91 habitual (compulsive) drug taking. As midbrain dopaminergic circuits are involved
92 in the establishment of habit learning (Balleine and Dickinson, 1998), this raises
93 the possibility that developmental ethanol exposure leads to an alteration in the
94 learning processes such that there is an increased propensity to develop habits,
95 and thus an increased likelihood of developing addiction.

96 During instrumental learning, responses are controlled by two distinct
97 processes. The first, response-outcome (R-O) learning, is seen early in the
98 learning process and is flexible in the sense that it is affected by alterations in
99 contingencies and by devaluation of the reinforcer. The second process,
100 stimulus-response (S-R), or 'habit' learning is far less flexible, is not affected by
101 alterations in contingencies or devaluation, and is often seen as a result of over-
102 training (Balleine and Dickinson, 1998). This form of behavioural plasticity has not
103 only been characterized in a variety of vertebrates (rodents; (Balleine and
104 Dickinson, 1998), primates; (Fernandez-Ruiz et al., 2001), horses; (Parker et al.,
105 2009)), but also in invertebrates (drosophila; (Brembs, 2009)). The shift between
106 R-O and S-R learning is adaptive: in a changing environment, the cost associated
107 with learning about potentially important new information must be balanced with
108 the ability to learn rules relating to continuously performed tasks. For example, it
109 would make driving a car a far more arduous task if we had to consult a manual
110 before every trip. In mammals these processes, collectively known as action
111 selection, are directly mediated by dopamine signalling between the prefrontal
112 cortex (PFC) and the dorsal striatum (DS). Specifically, the medial aspect of the
113 DS (DMS) is essential for R-O learning and the lateral aspect (DLS) for S-R
114 learning (Balleine and Dickinson, 1998).

115 Moderate prenatal ethanol exposure causes alterations in dendritic
116 morphology in the shell region of the nucleus accumbens (NAcs) in rats (Rice et
117 al., 2012b). The nucleus accumbens (NAc) is functionally and structurally
118 separated into the NAcs and the core region (NAcc). NAcs is primarily thought to
119 be involved with the processing of the incentive value of reward, whereas the
120 NAcc is controls instrumental learning (Corbit et al., 2001). Reductions in spine

121 density and dendritic length in the NAcS following prenatal ethanol exposure in
122 rats (Rice et al., 2012b) suggests that the increase in preference for ethanol is not
123 the result of a generalized increase in the incentive value of rewards. Instead, it
124 may be mediated by an augmented role for the NAcc in instrumental learning
125 increasing DS control of behaviour. In turn, this could lead to accelerated shift
126 from DMS-controlled R-O learning to DLS-controlled S-R habit (Yin et al., 2008).
127 We therefore designed this study to test the hypothesis that exposure to
128 moderate amounts of ethanol during early brain development would increase the
129 development of S-R habits in adulthood, using zebrafish as a model species.

130 Over recent years, the zebrafish has emerged as one of the major
131 vertebrate species in behavioural neuroscience, with procedures emerging
132 showing clear translational relevance to human disorders such as drug addiction
133 (Guo, 2004; Kalueff et al., 2013; Parker et al., 2013a). This has included
134 conditioned place preference for drugs of abuse (CPP; Brennan et al., 2011;
135 Darland and Dowling, 2001; Kily et al., 2008; Ninkovic et al., 2006), impulsivity
136 (Parker et al., 2014; Parker et al., 2013c) as well as compulsive drug seeking and
137 relapse (Brennan et al., 2011; Kily et al., 2008). **One of the key benefits of**
138 **using zebrafish to complement mammalian models is their small size and**
139 **genetic tractability. In addition, their transparency as embryos facilitates**
140 **exploration of developmental effects through *in vivo* imaging. This, coupled**
141 **with the fact that a number of brain systems linked to addiction in mammals**
142 **are well conserved in zebrafish, or at the very least, functional homologues**
143 **have been identified, makes them a potentially useful model. For example,**
144 **pallial structures such as the amygdala, hippocampus and piriform cortex**
145 **are conserved, and the ascending midbrain dopamine pathways (in**

146 mammals, the ventral tegmental area [VTA] to NAc to prefrontal cortex
147 [PFC] is present in teleost fish such as zebrafish (in zebrafish, the VTA
148 [posterior tuberal nucleus; PTN] and NAc [ventral [Vv] and dorsal [Vd]
149 telencephalic nuclei] (see Parker et al., 2013a for recent reviews; Parker and
150 Brennan, 2012). In addition, we have recently introduced zebrafish as a model
151 for moderate prenatal ethanol exposure, showing that adults exposed to low-
152 moderate levels of ethanol during early brain development have differences in
153 social cohesion and social exploratory behaviour and that these differences may
154 be mediated by differences in 5-HT circuits (Parker et al., In Press).

155 **Previous research in adult zebrafish found brain concentrations of**
156 **ethanol following chronic exposure typically reach ~80-90% bath**
157 **concentration in adults (Dlugos and Rabin, 2003; Mathur et al., 2011) and**
158 **~30-40% in embryos and larvae (Reimers et al., 2004). The relationship**
159 **between brain and blood alcohol content is not straightforward, and**
160 **estimates for the ratio of brain: blood ethanol range from 0.6-1.5 (Moore et**
161 **al., 1997). Therefore, we exposed embryos to 20mM ethanol which would**
162 **result in BAC of ~0.02-0.07g/dL, putting the zebrafish model in the moderate**
163 **prenatal ethanol classification (Valenzuela et al., 2012). The aims of this study**
164 were to test the hypotheses that moderate developmental ethanol exposure
165 would increase both drug preference and the formation of S-R habits. In addition,
166 we aimed to characterize mRNA expression in a variety of genes that encode
167 proteins related to addiction and habit formation, including the dopamine
168 receptors [DR] (Robbins and Everitt, 1999), dopamine transporter (DAT)
169 (Hironaka et al., 2004), nicotinic acetylcholine receptors [nAChRs] (Kitabatake et
170 al., 2003), μ -opioid receptor [MOR] (Miura et al., 2008), adenosine a2a receptor

171 (Yu et al., 2009), and noradrenaline transporter [NAT] (Xu et al., 2000). **We**
172 **chose these genes specifically as they have previously been shown to**
173 **affect aspects of habit formation and compulsive drug seeking. Many other**
174 **genes may also be involved with these behaviours and this is not meant as**
175 **an exhaustive list.**

For Review Only

176 **Materials and Methods**

177 *Subjects*

178 Adult Tubingen (mixed male/female) zebrafish were kept in a re-circulating
179 system, on a 14/10-hour light/dark cycle, at ~28.5°C within the zebrafish aquarium
180 facility at Queen Mary University of London. **There have been mammalian**
181 **studies that have distinguished between male and females. However, as the**
182 **genetic work was carried out on larvae, we would not be able to assign a sex to**
183 **the individuals at this age. Therefore, all animals were randomly assigned to**
184 **the treatment conditions.** Fish were fed with a mixture of flake food, fresh brine
185 shrimp and bloodworm. Adults were bred in house and fry reared according to
186 standard protocols (Westerfield, 1993). Larvae from each condition (20mM
187 ethanol and control) were sacrificed at 24hpf (i.e., before ethanol exposure),
188 50hpf (acute ethanol exposure: 2hrs after ethanol added), 72hpf and finally at
189 7dpf. All animal work was carried out following approval from the Queen Mary
190 Research Ethics Committee, and under licence from the Animals (Scientific
191 Procedures) Act 1986. Care was taken to minimize the numbers of animals used
192 in this experiment in accordance with the ARRIVE guidelines
193 (<http://www.nc3rs.org.uk/page.asp?id=1357>). Specifically, we examined data from
194 pilot studies in our lab, our previous published work (Parker et al., In Press) and
195 studies with other species (Rice et al., 2012b) to carry out a power calculation
196 and assess the minimum number of animals necessary for the expected effect
197 size with power of 0.8.

198

199 *Developmental ethanol exposure*

200

201 The developmental ethanol exposure is described in detail elsewhere (Parker et
202 al., In Press). Tubingen zebrafish larvae were treated by transferring them into
203 20mM (0.12% v/v, equating to ~0.04g/dL BAC) ethanol in aquarium water at 48
204 hpf. Prior to ethanol treatment, all larvae were staged to ensure they were at the
205 correct developmental stage (Kimmel et al., 1995). In addition, any larvae that
206 were still in their chorions were dechorionated to ensure equal exposure to
207 ethanol. We chose 48hpf to start treating the larvae as it represents a key stage
208 in CNS development, the Long Pec stage (Guo et al., 1999; Kimmel et al., 1995).
209 Furthermore, we quantified the size of larvae to ensure that there were no
210 differences between the treatment groups that might affect gene expression
211 analysis using a commercially available automated visualisation system
212 (www.elaborant.com). Previous research in adult zebrafish found brain
213 concentrations of ethanol following chronic exposure typically reach ~80-90%
214 bath concentration in adults (Dlugos and Rabin, 2003; Mathur et al., 2011) and
215 ~30-40% in embryos and larvae (Reimers et al., 2004). The relationship between
216 brain and blood alcohol content is not straightforward, and estimates for the ratio
217 of brain:blood ethanol range from 0.6-1.5 (Moore et al., 1997). Therefore, we
218 estimate that the larvae would have had a blood alcohol concentration (BAC) of
219 ~0.02-0.07g/dL putting the zebrafish model in the moderate prenatal ethanol
220 classification (Valenzuela et al., 2012).

221 Fish water was changed on alternate days, and the tanks were cleaned, in
222 order to reduce the build-up of yeast and control variation in the ethanol
223 concentration due to evaporation. Larvae were kept in Petri dishes until they were
224 five days old, after which they were transferred into tanks with dimensions 10 x 10
225 x 20 cm (depth x width x length cm) and a volume of 500ml with airlines. A

226 maximum of 50 fish were kept in each tank and numbers of ethanol and control
227 fish were balanced; dividing equally the quantity of fish receiving ethanol or
228 aquarium water for the controls between the tanks. Feeding of the larvae ZM000
229 and paramecium commenced at five days. After seven days swimming in fish
230 water containing ethanol (i.e., aged 9 days), the fry were transferred back to pure
231 aquarium water. At age three weeks the volume of water in the tanks was
232 increased to 1L to provide more space for growth and the ZM000 was replaced
233 with ZM100 and artemia. At age five weeks they were transferred into 7L tanks
234 and cleaning was reduced to once a week. Control fish were treated identically,
235 except that the ethanol was not added to their water at any time.

236

237 *Conditioned place preference*

238

239

240 The conditioned place preference procedure comprised an opaque tank
241 measuring (W x L x H) 16.5cm x 33cm x 15 cm and containing 3L of water (see
242 Parker et al., 2013b for detailed methods). The visual discriminanda (black spots
243 or black vertical lines) were located on the walls of the tanks (see supplementary
244 Figure 1). Conditioning started on a Monday, when the fish were habituated. All
245 fish were first placed individually in the conditioning tanks, with no ethanol added,
246 for a period of 20-mins. This was repeated on the second day (Tuesday), during
247 which time baseline preference for spots or stripes was recorded by measuring
248 time spent in each area of the tank. On the following two days (Wednesday-
249 Thursday), the fish were conditioned to their least-preferred stimulus, as
250 determined by their baseline preference. During conditioning trials, all fish were
251 initially exposed to the non-drug side for a period of 20 minutes. An opaque

252 barrier in the tank prevented access to the drug side. After 20-minutes, the barrier
253 was lifted to allow the fish to move to the drug-paired side of the tank. On entry,
254 the barrier was replaced and ethanol was added to the tank water. **In order to**
255 **avoid a concentration gradient through the tank, we carefully poured half of**
256 **the ethanol solution into opposite corners at the far ends the tank (see**
257 **supplementary Figure 1). We added pure ethanol mixed with distilled H₂O to**
258 **the tanks to avoid 100% ethanol being poured directly onto the fish, which**
259 **may have been aversive. In other words, 1.5 ml of ethanol was added to**
260 **48.5ml aquarium water, and this solution was added to the test tank.** This
261 procedure was repeated on the following 2 days. Following conditioning, all fish
262 were tested in a single probe trial, during which they were placed in a tank with
263 the barrier lifted such that both sides could be accessed freely. Time spent in the
264 proximity to the drug-paired stimulus was recorded. **This procedure was used**
265 **initially to determine the appropriate dose of ethanol. To do this, we tested**
266 **control fish at 0, 0.5, 1, and 1.5% ethanol. As a result 0.5% ethanol was**
267 **chosen as a threshold dose to use when comparing responses of**
268 **developmentally treated and control fish.**

269

270 *Habit formation*

271

272 **In order to test habit formation, we used a fish-version of the widely used**
273 **Tolman T-maze assay (supplementary Figure 2). This is the first time this**
274 **has been used in fish, but has been used in mammals regularly to measure**
275 **aspects of habit formation and S-R learning (Packard and Knowlton, 2002;**
276 **Yin and Knowlton, 2004; Yin et al., 2004).** Fish were initially habituated to the
277 apparatus for two half-hour sessions prior to testing. During training sessions, the

278 fish was initially placed in the test area in the South arm (nb, the release arm and
279 choice arms were counterbalanced between fish). The barrier was then removed
280 to allow the fish access to the choice arms. The barrier was closed behind the
281 fish. If the fish chose the East arm, a barrier was placed behind it and it was
282 reinforced with a small amount of brine shrimp (correct response). The barrier
283 remained in place for 20-seconds to ensure the fish had time to consume the
284 food. If the fish entered the West arm, it was given 60 seconds to choose the
285 correct arm, after which it was guided into the correct arm and fed. This ensured
286 that each trial was reinforced to remove any potential differences in reinforcement
287 history between the fish. Each fish was trained for 5 trials at a time (one training
288 block = 5 trials) until they reached criterion (< 20 -sec mean approach latency for 5
289 consecutive sessions and mean correct responses $\geq 75\%$). Once the fish had
290 reached criterion, they were tested in a single probe trial. This was always carried
291 out in the fifth trial of a 5-trial block. In other words, trials 1-4 were normal training
292 trials, and trial 5 was the probe. During the probe trial everything was exactly the
293 same as during training trials except that the fish was released from the opposite
294 arm to that from which it was released in training. During the probe trial, no
295 reinforcement was given. The experimenter noted which arm the fish entered on
296 release from the starting arm. The experimenter was blind to treatment allocation
297 at all times during testing to eliminate any bias (fish were allocated a number, and
298 their group identity not revealed before data analysis).

299

300 *qPCR procedures*

301

302 Larvae at 24hpf, 50hpf, 72hpf and 7dpf in batches of $n = 3$ embryos were

303 homogenized in 200 μ l Lysis buffer with 2 μ l Proteinase K for 30-45min (55°C).

304 mRNA was isolated using 80µl Dynabeads® Oligo(dT)₂₅ according to
305 manufacturer's instructions. Reference genes (see Table 1 for primer sequences)
306 were chosen according to previous research *β-actin*, *ef1a* and *rpl13a* (Parker et
307 al., In Press; Tang et al., 2007). Target genes used were nicotinic acetylcholine
308 receptors (nAChRs; *chrna2-7*, *chrnb2-4*), dopamine receptors and transporter
309 (DAT; *drd1-3*, *slc6a3*), noradrenaline receptor (NAT; *slc6a2*) and µ-opioid
310 receptor (MOR; *oprm1*). Absolute quantification was obtained by making
311 standards for each gene, prepared using the relevant primers to amplify
312 fragments from cDNA. Samples were then PCR purified and diluted to 10¹¹
313 fragments using the Avogadro constant. All qPCR reactions were carried out in
314 triplicate. 2µl of cDNA and 2µl each of forward and reverse primers (see Table 1)
315 was added to 10µl SYBR® Green PCR Master mix (Applied Biosystems) on a
316 LightCycler LC480 instrument (Roche Diagnostic). For detailed methods see
317 (Gemenetzidis et al., 2010; Teh et al., 2013).

318 [TABLE 1 HERE]

319 *Data analysis*

320

321 Statistical analyses were carried out in IBM SPSS for Macintosh (version 21). All
322 null hypotheses were evaluated with respect to $\alpha = 0.05$. Normality,
323 homoscedacity and homogeneity of variance were ascertained by examination of
324 fitted vs. residual plots from the models. **With respect to the CPP dose-**
325 **response curve, data were fitted to a quadratic regression (dependent**
326 **variable = change in preference for the drug-predictive cue from baseline,**
327 **independent variable = dose of ethanol (0, 0.5, 1, 1.5% v/v).** Data pertaining to
328 the comparison of CPP between ethanol-treated and control fish was analysed
329 using a 2-way mixed design analysis of variance (ANOVA), with one between-

330 subjects factor (group: ethanol-exposed vs. handling control), one within-subjects
331 factor (phase: pre- vs. post-conditioning), as well as their interaction (group x
332 phase). The dependent variable was time spent in proximity to the conditioned
333 cue.

334 With respect to T-maze acquisition, the effects of developmental ethanol
335 treatment on latency to approach the target arm, and proportion of correct
336 responses, were fitted by restricted maximum likelihood (REML) to linear mixed
337 effects model (LME). Denominator degrees of freedom were estimated using the
338 Satterthwaite approximation (Satterthwaite, 1946). Latency data were fitted to a
339 model with Gaussian error structure by REML, and correct responses were fitted
340 to a logistic regression model (binomial distribution) by Laplace approximation.
341 For the binomial model, significance was ascertained from Type-III Wald Chi-
342 Squared statistics performed on the final model. Training block (1-31), group
343 (ethanol-exposed vs. handling control) and their interaction, were added to initial
344 models as fixed effects. Fish-ID was added to both models as a random effect to
345 control for associated interclass correlation (random-intercept model). Finally,
346 probe trial data (frequency of 'habit' vs. 'place' responses for ethanol-exposed or
347 handling control fish) was analysed using a Fisher's exact test. qPCR data were
348 analysed with 2-way factorial ANOVAs. The between-subjects factors were age
349 (4-levels: 24hpf, 48hpf, 72hpf, 7dpf) and treatment (2-levels: ethanol vs handling
350 control). Descriptive statistics are reported as mean \pm SE.

351 **Results**

352

353 ***Moderate ethanol exposure during early brain development does not affect***354 ***unconditioned locomotor behaviour***

355

356 **First, we examined the basal locomotor responses of the adult fish that had**357 **been exposed as embryos to 20mM ethanol. This information is extremely**358 **relevant, since ethanol may impair motor coordination depending on both**359 **time of exposure and concentration. Without this, it may be difficult to**360 **comprehend how such data can be related to memory, since behavioural**361 **tasks in animal models is directly dependent of their locomotor profile. We**362 **placed the fish individually (n = 14 treated with 20mM ethanol as embryos, n**363 **= 14 control) in a circular container (radius = 30cm) and filmed them for 10**364 **min (Noldus Ethovision, TrackSys, Nottingham UK). We measured the total**365 **distance travelled over the 10 mins (cm) as well as the number of rotations**366 **(i.e., complete body rotations about the central axis of the body) to examine**367 **any gross motor effects of the ethanol. Independent t-tests revealed no**368 **effects of ethanol on distance ($t [26] = 1.09, P = 0.29$; control = 9427 ± 654.02** 369 **cm 20mM ethanol = 8475.38 ± 581.04 cm) or rotations ($t [26] = 0.39, P = 0.7$;**370 **control = 16.29 ± 4.2 rotations, 20mM ethanol = 19.43 ± 6.97 rotations).**

371

372 ***Moderate developmental ethanol exposure increases CPP for ethanol***373 **[FIGURE 1 ABOUT HERE]**

374

375

376 **Figure 1A displays the dose-response curve to ethanol. There was a**
377 **significant quadratic effect on change in preference as a function of dose,**
378 **$F(2, 42) = 7.89, P = 0.001, R^2 = 0.27$ (regression parameter estimates: $y = -$**
379 **$0.04+0.41*x^2-0.2*x$). The highest point on the peak was at 1%, but to avoid**
380 **potential ceiling effects in the developmental ethanol experiment, we used**
381 **0.5% v/v to condition our fish.**

382 Figure 1B displays the time spent in the vicinity of the conditioned cue (i.e.,
383 spots or stripes, depending on least preferred during basal preference
384 assessment) before (basal) and after (probe trial) conditioning with 0.5% ethanol.
385 Fish that were developmentally exposed to 20mM ethanol showed a greater
386 increase in preference for the drug-paired cue following conditioning. These
387 patterns were confirmed with an LME. There was a main effect of phase, $F(1,75)$
388 $= 16.28, p < 0.001$, but not group, $F < 1$. There was also a significant group x
389 phase interaction, $F(1,75) = 5.3, p = 0.02$, characterized as the ethanol exposed
390 group showing a greater increase in preference for the conditioned cue during the
391 probe trial ($p < 0.01$). In summary, developmental exposure to 20mM ethanol
392 significantly augmented the increase in preference for the drug-paired cue
393 following CPP training with 0.5% v/v ethanol.

394

395 *Moderate developmental ethanol exposure increases motor habit formation*

396 [FIGURE 2 ABOUT HERE]

397

398 Figure 2 displays the acquisition data for fish during training on the T-maze
399 assay. Performance was variable throughout in both groups, but ethanol treated
400 fish showed marginally poorer learning during the training overall. In terms of
401 latency to respond (Figure 2A), an LME revealed no main effect of group, $F < 1$,

402 but there was a main effect of block, $F(30, 1788) = 2.43, p < 0.0001$ and a
403 significant group x block interaction, $F(30, 1788) = 1.79, p < 0.01$. The interaction
404 was characterized as the ethanol fish showing longer latencies in blocks 3, 18, 21
405 and 22 ($ps < 0.05$).

406 In terms of correct responses during each session, we fitted a logistic
407 regression model by Laplace approximation. There was a significant main effect
408 of block (Figure 2B), $\chi^2(30) = 103.26, p < 0.0001$, with correct responses
409 increasing after block 21. There was a marginally non-significant main effect of
410 group, $\chi^2(1) = 3.37, p = 0.06$, and there no significant group x block interaction,
411 $\chi^2(30) = 33.9, p = 0.28$. To summarise, all fish appeared to have learnt the
412 discrimination, although there was some evidence from latency data that the fish
413 exposed developmentally to 20mM ethanol learnt at a slightly slower rate than
414 handling controls.

415 During the probe trial, 75% of the fish exposed to ethanol appeared to
416 show a motor habit, turning in the direction they had turned during training. Of the
417 fish from the handling control group, 100% went to the arm reinforced during
418 training in the probe trial, suggesting that none of them had adopted a habit. This
419 difference was confirmed with a Fisher's exact test (Figure 2C; $p = 0.01$).

420

421 *Moderate developmental ethanol exposure affects mRNA expression of nicotinic,*
422 *dopaminergic and μ -opioid receptors*

423 [FIGURE 3 ABOUT HERE]

424

425 There was an effect of ethanol treatment on *chrna2* mRNA expression that
426 approached significance, $F(1,24) = 3.96, P = 0.058$, and a significant effect of

427 age, $F(3,24) = 6.4$, $P < 0.01$. There was also a significant age x treatment
428 interaction (see Figure 3A), $F(3,24) = 3.92$, $P < 0.05$, characterized as
429 expression being significantly suppressed in ethanol-treated 50hpf embryos [$P <$
430 0.05]. There was a significant effect of age on *chrna3* mRNA expression, $F(3,24)$
431 $= 5.38$, $P < 0.01$, but no significant effect of prenatal ethanol treatment, $F(1,24) =$
432 2.12 , $P = 0.16$. There was, however, a significant age x treatment interaction (see
433 Figure 3B), $F(3,24) = 2.95$, $P = 0.05$, characterized as expression being
434 suppressed at 50hpf in the ethanol-treated embryos [$P < 0.05$]. There was a
435 significant effect of age on *chrnb3* expression, $F(3,24) = 14.68$, $P < 0.01$, but no
436 effect of treatment, $F(1,24) = 1.57$, $P = 0.22$. There was also a significant age x
437 treatment interaction (see Figure 3C), $F(3,24) = 7.45$, $P = 0.001$, characterized
438 as ethanol causing an increase in mRNA expression at 50hpf [$P < 0.01$], followed
439 by a reduction in expression at 7dpf [$P < 0.05$].

440 There was a significant effect of age on *slc6a3* mRNA expression, $F(3,$
441 $24) = 5.67$, $P < 0.01$, but no effect of treatment, $F(1,24) = 3.67$, $P = 0.07$. There
442 was a significant age x treatment interaction (see Figure 3D), $F(3,24) = 3.45$, $P =$
443 0.03 characterized as there being an increase in mRNA expression at 50hpf, but
444 only in the control group (i.e., expression was suppressed by ethanol treatment) [P
445 $= 0.03$]. There was a significant effect of age on *drd3* mRNA expression, $F(3,24)$
446 $= 5.55$, $P < 0.01$, but no effect of treatment, $F < 1$. There was a significant age x
447 treatment interaction (see Figure 3E), $F(3,24) = 3.27$, $P = 0.04$ characterized by
448 expression being significantly increased in the ethanol-treated group at 50hpf [$P <$
449 0.01], but this effect reversed at 7dpf [$P < 0.05$]. There was a significant effect of
450 age on *oprm1* mRNA expression, $F(3,24) = 4.55$, $P = 0.012$, but no effect of
451 treatment, $F < 1$. There was also a significant age x treatment interaction (Figure

452 3F), $F(3,24) = 3.26$, $P < 0.05$, characterized as ethanol significantly increasing
453 mRNA expression at 50hpf [$P < 0.05$] but this effect reversed at 7dpf [$P < 0.05$].

454 There were no significant effects of ethanol treatment on mRNA
455 expression for the other nAChR-encoding genes (*chrna4*, *chrna5*, *chrna6*,
456 *chrna7*, *chrnb2*, *chrnb4*), nor those encoding for the other dopamine receptors
457 (*drd1*, *drd2*), adenosine receptor (*adr2a*, *adr2b*) or noradrenaline transporter
458 (*slc6a2*), nor were there any age x treatment interactions (see **Supplementary**
459 **Results**).

460

461

462 Discussion

463 Exposure to moderate levels of ethanol during early brain ontogeny
464 increases preference for alcohol and other drugs in the offspring, and acts as a
465 significant risk factor for the development of addiction. The aim of this experiment
466 was to explore the mechanism by which this manifests by examining behavioural
467 and gene expression changes following exposure to moderate levels of ethanol
468 during brain development in zebrafish. Exposure to 20mM (0.12% v/v, equivalent
469 to 0.2-0.7 g/dL BAC) ethanol from 48hpf-9dpf increased the time spent in the
470 presence of the ethanol-conditioned cue in the CPP test as compared to non-
471 exposed controls, suggesting an increased preference for alcohol in the exposed
472 fish as has been previously observed in rodents. **We also found that after**
473 **training on a place-response test in a modified Tolman T-maze, there was a**
474 **difference in performance at the specific time point at which the criterion as**
475 **selected were met, suggesting that exposed fish were more likely to show**
476 **S-R learning early in the learning process, whereas non-exposed controls**

477 **showed R-O learning.** This indicated an increased propensity for habit formation
478 in developmentally exposed individuals. Finally, we found that ethanol acutely
479 suppressed *chrna2*, *chrna3* and *slc6a3* mRNA expression. It also acutely
480 increased *chrnb3*, *drd3* and *oprm1* mRNA expression, but this effect reversed by
481 7dpf for all three suggesting ethanol caused adaptation in expression of these
482 genes during early brain development. Our data support the hypothesis that
483 animals developmentally exposed to ethanol show increased S-R learning and
484 suggests involvement of dopaminergic, opioid and nicotinic pathways in this
485 process.

486 We found that ethanol increases preference for alcohol in a CPP test, in
487 agreement with previous work showing that prenatal ethanol exposure increases
488 alcohol preference in later life in humans (Spear and Molina, 2005) and in rodents
489 (Arias and Chotro, 2005; Van Waes et al., 2011). Importantly, this shows the
490 translational relevance of fish in this procedure. Two main theories to explain
491 increased self-administration seen following developmental exposure to ethanol
492 exist in the literature. The first is that ethanol preference is caused by the
493 chemosensory adaptation of the foetus developing in an alcoholic medium
494 (Dominguez et al., 1993; Faas et al., 2000). This theory has been challenged in
495 particular by the observation that exposed offspring show a higher preference for
496 other rewarding drugs as well as ethanol (Barbier et al., 2009). An alternative
497 theory is that ethanol alters structural and functional neuroplasticity, particularly in
498 the striatum and other midbrain structures, and this alters reward circuitry. There
499 is some evidence for changes in neural pathways associated with reward
500 following developmental ethanol (Blanchard et al., 1993; Choong and Shen,
501 2004; Malanga and Kosofsky, 1999; Middaugh et al., 1994). However, an

502 increased preference for alcohol (or other drugs) does not necessarily translate in
503 to an increased propensity for addiction (Everitt et al., 2008). Indeed, preclinical
504 studies have found that there is no clear correlation between self-administration
505 rates and the development of compulsive drug seeking or relapse, suggesting
506 that other processes are more important (Belin et al., 2008).

507 In addition to an increased preference for ethanol, we found that fish
508 exposed to ethanol during early brain development appeared to shift from R-O to
509 S-R earlier in the learning process. Although they learned the task marginally
510 slower than the controls, in a probe trial where the fish were released from the
511 opposite side of the maze, the ethanol fish all showed a motor habit, whereas the
512 majority of the controls used the extra-maze cues to guide navigation. This
513 difference in stimulus control relatively early in the learning process suggests that
514 the ethanol exposed animals were displaying less goal-directed, cognitive control
515 over their behaviour, and more automated 'habitual' responses, an effect typical
516 of those addicted to drugs (Balleine and Dickinson, 1998). Thus, our findings
517 suggest that exposure to ethanol during early brain development enhances habit
518 formation, potentially rendering the exposed animal more susceptible to addiction.
519 This is particularly interesting in the light of the previously discussed brain
520 changes observed following developmental ethanol exposure in rodents, where
521 rats exposed to moderate ethanol during gestation had a decrease in MSN spine
522 density and dendritic length in the NAc (Rice et al., 2012b). As these regions are
523 key to the process of instrumental contingency learning, this provides a
524 mechanism potentially linking developmental ethanol to enhanced habit formation
525 (i.e., ventral to dorsal shift; Belin et al., 2013), fitting with current theories that
526 characterise addiction in terms of maladaptive habits where drug seeking comes

527 under the control of conditioned reinforcers (drug-related discriminative stimuli)
528 (e.g., Everitt and Robbins, 2005; Everitt and Robbins, 2013).

529 As developmental ethanol exposure has been shown to alter gene
530 expression in a number of pathways, we examined expression of components of
531 pathways associated with the switch from R-O to S-R learning (e.g., dopamine,
532 opioid and adenosine) as well as other systems associated with drug seeking and
533 addiction (adrenergic and cholinergic) in larval fish. We found that ethanol
534 caused a number of changes in mRNA expression in genes relevant to addiction.
535 First, we found that ethanol initially increased and then decreased, *drd3*
536 expression, the gene that encodes for the DRD3. This supports previous work
537 with rats that showed prenatal ethanol exposure reduced quiniprole (a DRD2/3
538 agonist)-induced yawning (Brus et al., 1995). Lower levels of DRD2/3 in the
539 ventral striatum have been linked both to poor inhibitory control and to escalation
540 of self-administration rates for cocaine in rodents (Dalley et al., 2007). In addition,
541 DRD3 expression has been linked specifically to habit formation, with
542 schizophrenia patients and healthy controls with a variant (Ser9Gly
543 polymorphism) at the DRD3 locus showing stronger habit learning (Kéri et al.,
544 2005). We did not see any differences for DRD1 or DRD2 expression, or in the
545 adenosine a2a receptor (which forms a heterodimer with DRD2; Franco et al.,
546 2000), but we did see an acute effect on DAT with ethanol-exposed embryos
547 showing suppression of *slc6a3* mRNA expression at 50hpf (2hrs after ethanol
548 was added to the embryo medium), with expression normalising thereafter. DAT-
549 KO mice show resistance to extinction, suggesting increases in habit-like learning
550 (Hironaka et al., 2004). Here, there were no differences in expression at 7dpf, but
551 we did not test the fish in adulthood for DAT. The fact that *slc6a3* mRNA

552 expression was acutely suppressed by ethanol (i.e., at 50hpf) suggests that there
553 may be some long-term effects on general DAT function, and this could be
554 studied in more detail in future experiments.

555 In addition to finding what appeared to be an adaptive change in DRD3,
556 we observed an acute increase in *orpm1* mRNA expression (50hpf) followed by a
557 decrease in expression at 7dpf. Again, this is suggestive of adaptation of
558 expression following initial change caused by the ethanol. This gene codes for
559 the μ -opioid receptor (MOR) protein. With respect to drug preference, MOR-KO
560 mice (homozygous and heterozygous) show reduced self-administration of
561 ethanol (Hall et al., 2001). MORs are widely distributed around the mammalian
562 ventral and dorsal striatum (Arvidsson et al., 1995) and have been linked to the
563 appetitive phase of instrumental learning (Mucha and Herz, 1985). This suggests
564 that a reduction in MOR will reduce the motivational value of rewards, thus
565 rendering the animal less sensitive to Pavlovian appetitive cues or conditioned
566 reinforcers (Balleine and Dickinson, 1998). This is somewhat at odds with our
567 observations here that ethanol-exposed fish show a reduction in MOR mRNA
568 expression at 7dpf, but an increased preference for ethanol in the CPP test.
569 Theories linking MOR genetic variants to addiction are common (Contet et al.,
570 2004). Depletion of MOR during learning by naloxone causes an increase in
571 habit-based learning (Wassum et al., 2009) consistent with our findings here. As
572 such, despite this mechanism being fairly speculative at this stage, it certainly
573 warrants further investigation.

574 In addition to changes in DRD3, DAT and opioid receptor we observed that
575 the ethanol-exposed larvae had differences in nAChR mRNA expression.
576 Specifically, we found an initial increase in *chrb3* expression after acute ethanol

577 exposure (50hpf), followed by a decrease in expression at 7dpf. We also saw
578 suppression of both *chrna2* and *chrna3* mRNA expression following acute
579 exposure, but no differences in expression in these genes later in development
580 (i.e., at 7dpf). *chnrb3* encodes the β -3 nAChR subunit (nAChR-b3). Humans with
581 specific allelic variants at this locus have previously been shown to show stronger
582 preference for nicotine (Zeiger et al., 2008) and alcohol (Hoft et al., 2009). This
583 latter study may explain why our fish showed a stronger preference for ethanol in
584 the CPP test. However, as stated earlier, increased preference for drugs does not
585 necessarily translate into, nor is it necessarily a predictor for, addiction.
586 Interestingly, and of relevance to our observation that ethanol exposure increases
587 habit-learning, variants at the nAChR-b3 locus have recently been shown to
588 predict differences in addictive behaviour, i.e., as measured by the Fagerström
589 Test for Cigarette Dependence (Rice et al., 2012a). This suggests that variation
590 at this locus may predict behaviours that are related to dependence rather than
591 simply heavy use of drugs (e.g., compulsive drug use). In mammals, nAChRb3s
592 are located in a the form of hetero-oligomer populations ($\alpha 6\beta 2\beta 3$, $\alpha 6\alpha 4\beta 2\beta 3$), on
593 DS dopamine neurons the region of the striatum responsible for the shift from
594 cognitive to habit learning (Balleine and Dickinson, 1998). They are also located
595 on VTA dopamine neurons ($\alpha 6\beta 2\beta 3$), a key component of the mesoaccumbens
596 dopamine pathway, and on neurons in the habenula-interpeduncular (IPN)
597 system ($\alpha 3\beta 3\beta 4$), thought to mediate aspects of withdrawal from drugs
598 (Changeux, 2010; Grady et al., 2009). In zebrafish, the same nAChR subunits
599 have been identified and serve a similar function as in mammals; they are located
600 mainly pre-synaptically and mediate the release of other neurotransmitters (e.g.,
601 dopamine) (Ackerman et al., 2009; Zirger et al., 2003). Therefore, nAChR-b3 may

602 be important for the regulation of dopamine release in the striatum during learning
603 (Azam et al., 2002; Cui et al., 2003; Zhou et al., 2001), thus helping to facilitate
604 the process of habit learning.

605 Finally, we found that ethanol was related to changes in *chrna2* and
606 *chrna3* mRNA expression following acute ethanol administration (50hpf). These
607 changes were not observed later in development. *chrna2*, which encodes for the
608 $\alpha 2$ nAChR subtype (nAChRa2) and *chrna3*, which encodes for the $\alpha 3$ nAChR
609 (nAChRa3) subtype, are expressed in the zebrafish IPN and habenula,
610 respectively, in a similar fashion to mammals (Hong et al., 2013). nAChRa2 in the
611 IPN (Salas et al., 2009) and nAChRa3 in the habenula (Jackson et al., 2013)
612 have been linked to the withdrawal from nicotine in rodent models, but not yet to
613 withdrawal from other drugs of abuse, such as alcohol. However, both the IPN
614 and the habenula receive major afferent inputs from the hypothalamus, limbic
615 forebrain and mesoaccumbens pathway (Bianco and Wilson, 2009), and may
616 regulate dopamine activity in reward pathways (Matsumoto and Hikosaka, 2007).

617 **Finally, in this study we chose to use embryos at different**
618 **developmental stages to examine gene expression. The reason for this was**
619 **to exclude the possibility of social or environmental variables (which we did**
620 **not vary systematically in this study) affecting gene expression. We accept**
621 **that there is a possibility that phenotypes will alter as a result of gene-**
622 **environment interactions during development, and the precise roles that**
623 **this has to play in the development of different behavioural phenotypes**
624 **(e.g., addiction) in later life would be an interesting area of future study.**

625 In summary, we tested the hypotheses that exposure to moderate levels of
626 ethanol during early brain ontogeny would result in enhanced place preference

627 for ethanol and increased habit formation during learning. **We found support for**
628 **both of these hypotheses, indicating that in fish, exposure to ethanol**
629 **developmentally has an impact on later behaviour and learning.** We also
630 found that ethanol exposure during this critical developmental stage caused
631 specific changes in dopamine and nicotinic receptor gene expression.
632 **Collectively, these data suggest a possible molecular mechanism by which**
633 **prenatal alcohol exposure may increase the risk of addiction in later life.**
634 **The translational relevance of this in the fish is borne out in the structural**
635 **and functional homology in brain regions related to these behavioural**
636 **systems and neural pathways between the species (Guo, 2004; Parker et al.,**
637 **2013a; Parker and Brennan, 2012), and existing evidence that humans**
638 **exposed to moderate levels of ethanol prenatally are susceptible to**
639 **addiction in later life (Baer et al., 2003).** Although speculative at present, it
640 may be that ethanol exposure during early brain development causes subtle
641 changes in the balance of nAChR in the IPN-habenular pathway that affects
642 development of dopamine systems. This model will need to be examined in more
643 detail in the future, for example by exposing both nAChR or DAT KO animals to
644 ethanol and examining mRNA expression and behaviour during development and
645 in adulthood.

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Figure 1. Time spent in vicinity of conditioned cue after 2 conditioning sessions in CPP task for fish developmentally exposed to 20mM ethanol and handling controls. ** $p = 0.01$.

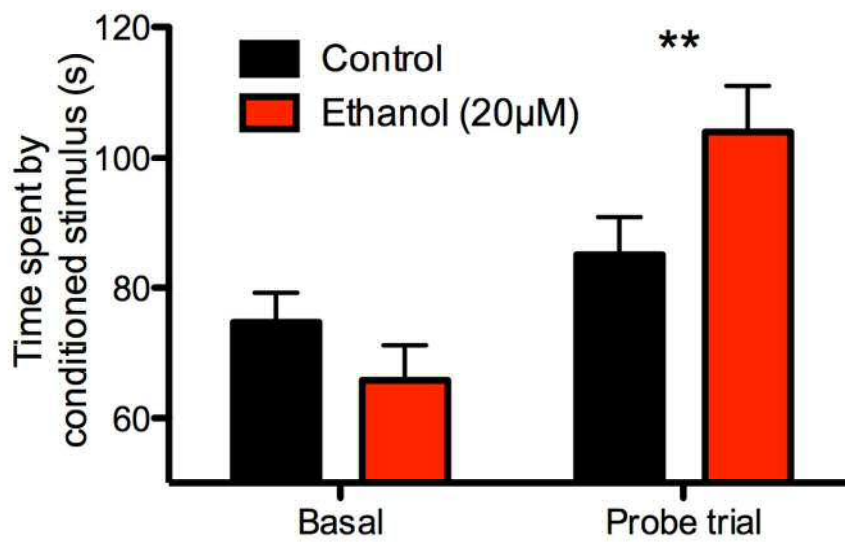
Figure 2. Acquisition graphs for training on T-maze. Left hand panel displays the proportion of correct trials in each session (A); Right hand panel displays the approach latency (s) (B);. Learning criterion was set at both groups showing mean approach latencies of < 20 sec for 5-consecutive trials and both groups >75% correct.

Figure 3. Changes in mRNA expression of nAChRs *chrna2* (A), *chrna3* (B), *chrnb3* (C), dopamine transporter (D) and dopamine D3 receptor, *drd3* (E), and μ -opioid receptor, *oprm1* (F), normalised against β -actin, *ef1a* and *rpl13a*, following exposure to moderate levels of ethanol during early brain development.

Supplementary Figure 1. Arena used for CPP assay. Basal preference was ascertained, and the fish was conditioned to the least preferred side. Drug was added to both sides of the apparatus to ensure equal dispersion across the tank.

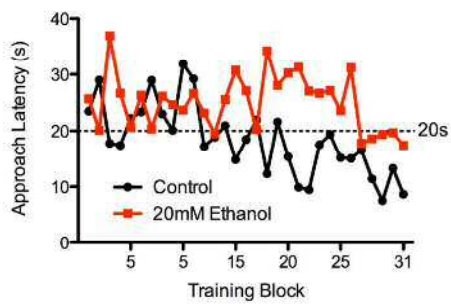
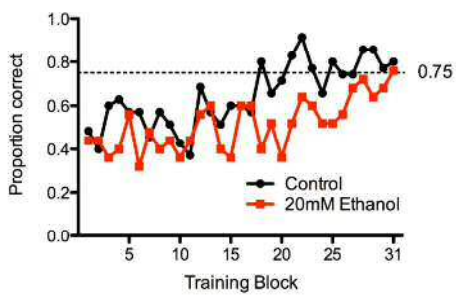
Supplementary Figure 2. T-maze used to test fish in a place-response test. Fish were initially entered into the south (S) arm (nb, this was counterbalanced between fish, so half started in the north (N) arm). The barrier was released and the fish was reinforced for entries to the east (E) arm (nb, this was counterbalanced, so half were reinforced in the west (W) arm). During the probe trial, the fish was released from the opposite arm to that from which it was released during training, and the direction of its response was noted.

Table 1. *Primer Sequences for Quantitative Real-Time PCR Assessment of mRNA Expression*



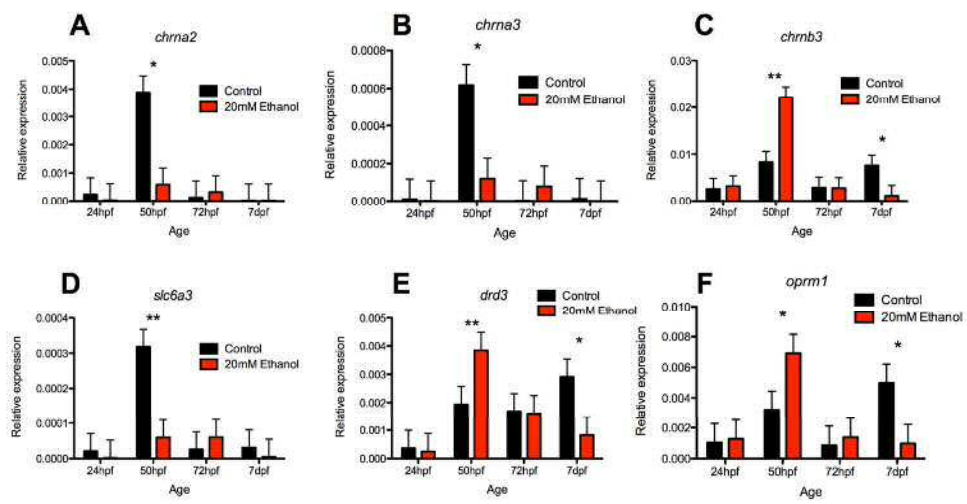
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View Only



249x90mm (300 x 300 DPI)

Review Only

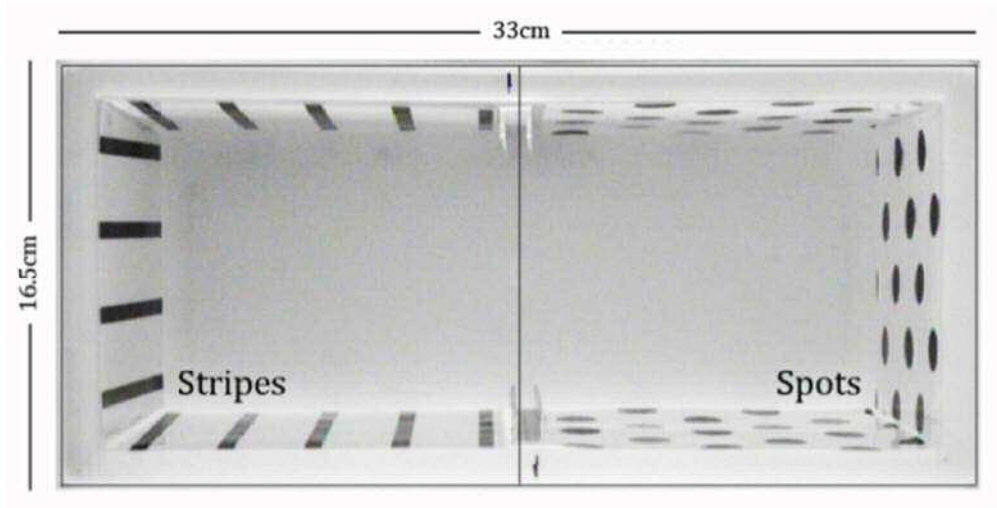


248x132mm (300 x 300 DPI)

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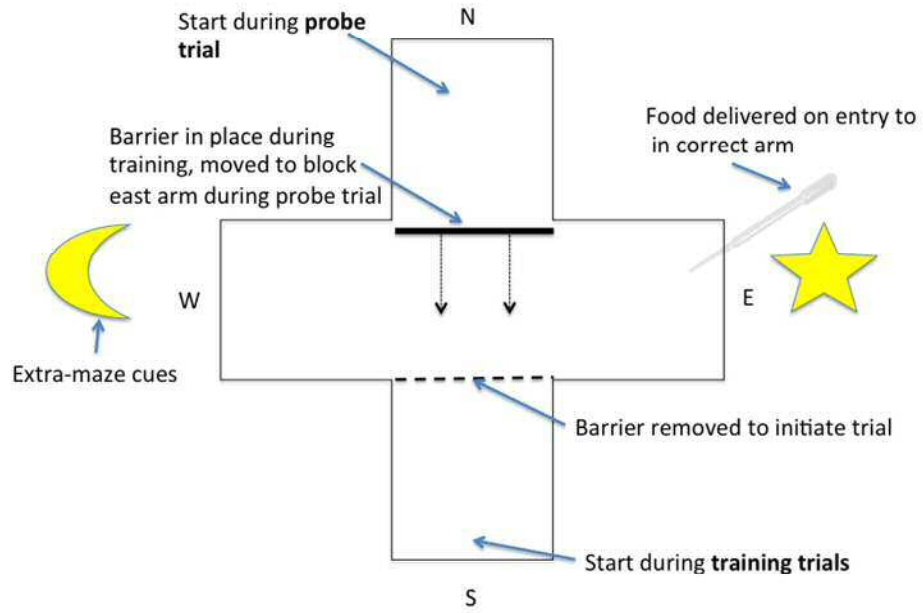
Gene name	Primers
<i>adora2aa-F</i>	CTT GAG CGC AGG AAC CAG AG
<i>adora2aa-R</i>	CGC GCA CTG AGA GAT GAC AG
<i>adora2ab-F</i>	CTG GTG ATT GCG GTT TTG GC
<i>adora2ab-R</i>	CAG GAC TCC GAC TGC AAT GT
<i>β-actin-F</i>	CGA GCT GTC TTC CCA TCC A
<i>β-actin-R</i>	TCA CCA ACG TAG CTG TCT TTC TG
<i>rpl13α-F</i>	TCT GGA GGA CTG TAA GAG GTA TGC
<i>rpl13α-R</i>	AGA CGC ACA ATC TTG AGA GCA G
<i>eF1α-F</i>	CTG GAG GCC AGC TCA AAC AT
<i>eF1α-R</i>	ATC AAG AAG AGT AGT ACC GCT AGC ATT AC
<i>chrna2-F</i>	GCG GAA AAC CGG ATA AAA ACA CTC
<i>chrna2-R</i>	AGT TTG TCC TCT GCG TGT GCA T
<i>chrna3-F</i>	TGT ACA TCC GCC GAT TAC CGC T
<i>chrna3-R</i>	TCC GCA GTC GGA GGG CAG TA
<i>chrna4-F</i>	TTA CAA GAG GTT TGG GCG CT
<i>chrna4-R</i>	ACA GAC CAG TAG ATC ATC ACT CC
<i>chrna5-F</i>	GGC TCC CAG GTC GAC ATT
<i>chrna5-R</i>	AAC CCC GGT TAC CAG TGG CCT
<i>chrna6-F</i>	CTT TGG GCC TCT TCC TGC AA
<i>chrna6-R</i>	TCA GAG TCT TGA TGT AGT GAC GG
<i>chrna7-F</i>	ACC GTG TCA CAT TGT TCA TTC TC
<i>chrna7-R</i>	ACA GGT CTC TCC AGT GGG TTA
<i>chrnb2-F</i>	GGC TGC CTG ATG TTG TTC TT

chrnb2-R TGG TGG CAA CCA GAA GAC ACT T
chrnb3-F CAG GAG TCA ACC TCC GCT TT
chrnb3-R TGA ATC TGA ACG CAC TGG CT
chrnb4-F TGA TCA CAT GAT GGG GAA TGA CG
chrnb4-R CAC CAC ACA CAC GAT CAC AAA G
drd1-F TGG TTC CTT TCT GCA ACC CA
drd1-R AGT GAT GAG TTC GCC CAA CC
drd2-F TCC ACA AAA TCA GGA AAA GCG T
drd2-R CAG CCA ATG TAA ACC GGC AA
drd3-F ATC GAG TTT CGC AGA GCC TT
drd3-R TCC ACA GTG TCT GAA AGC CG
oprm1-F CCG TAT GTG ACA GGA CGC CA
oprm1-R TTT CCC ACC AGT CCC ATC ACA
slc6a2-F AGG TGA CAT TGT TTG AGA TGT CTT
slc6a2-R TGT CTT GGT AGT GTC AAG TTG T
slc6a3-F TAT GTG GTC CTG ACC GTG CT
slc6a3-R CAC ATG TGT AGG CGC AGG AA



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